



1600

#9½ RSL

**RAW SEQUENCE LISTING**  
**PATENT APPLICATION: US/09/651,374A**

**DATE: 04/24/2002**  
**TIME: 09:14:19**

**Input Set : A:\A-68197.txt**  
**Output Set: N:\CRF3\04242002\I651374A.raw**

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3 <110> APPLICANT: Wittwer, Carl
4      Hermann, Mark
6 <120> TITLE OF INVENTION: Homogenous Multiplex Hybridization Analysis by Color and TM
8 <130> FILE REFERENCE: A-68197/RFT
10 <140> CURRENT APPLICATION NUMBER: US 09/651,374A
11 <141> CURRENT FILING DATE: 2000-08-30
13 <150> PRIOR APPLICATION NUMBER: US 60/151,494
14 <151> PRIOR FILING DATE: 1999-08-30
16 <160> NUMBER OF SEQ ID NOS: 30
18 <170> SOFTWARE: PatentIn version 3.1
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23 <213> ORGANISM: Homo sapiens
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65 <213> ORGANISM: Homo sapiens
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68 gcggctcctg cccgatgccg atgacctgca gaagtgcctg gccagtgtac ca	52
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74 <213> ORGANISM: Homo sapiens	
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77 gcggctcctg cccgatgccg atgacctgca gaagcgccctg gcagtgtacc a	51
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82 <212> TYPE: DNA	
83 <213> ORGANISM: Artificial Sequence	
85 <220> FEATURE:	
86 <223> OTHER INFORMATION: synthetic.	
88 <400> SEQUENCE: 7	
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93 <211> LENGTH: 27	
94 <212> TYPE: DNA	
95 <213> ORGANISM: Artificial Sequence	
97 <220> FEATURE:	
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109 <220> FEATURE:	
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121 <220> FEATURE:	
122 <223> OTHER INFORMATION: synthetic.	
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129 <211> LENGTH: 19	
130 <212> TYPE: DNA	
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133 <220> FEATURE:	
134 <223> OTHER INFORMATION: synthetic.	
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205 <220> FEATURE:  
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213 <211> LENGTH: 214  
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 222 gtggggcaag gtgaacgtgg atgaagttgg tggtgaggcc ctggcaggt tggtatcaag 180  
 224 gttacaagac aggttaagg agaccaatag aaac 214  
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 229 <212> TYPE: DNA  
 230 <213> ORGANISM: homo  
 232 <400> SEQUENCE: 19  
 233 agtcagggca gagccatcta ttgcttacat ttgcttctga cacaactgtg ttcactagca 60  
 235 acctcaaaca gacaccatgg tgcacctgac tcctgaggag aagtctgccg ttactgccct 120  
 237 gtggggcaag gtgaacgtgg atgaagttgg tggtaaggcc ctggcaggt tggtatcaag 180  
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 248 agtcagggca gagccatcta ttgcttacat ttgcttctga cacaactgtg ttcactagca 60  
 250 acctcaaaca gacaccatgg tgcacctgac tcctgaggag aagtctgccg ttactgccct 120  
 252 gtggggcaag gtgaacgtgg atgaagttgg tggtgaggcc ctggcaggt tggtatcaag 180  
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 265 acctcaaaca gacaccatgg tgcacctgac tcctaaggag aagtctgccg ttactgccct 120  
 267 gtggggcaag gtgaacgtgg atgaagttgg tggtgaggcc ctggcaggt tggtatcaag 180  
 269 gttacaagac aggttaagg agaccaatag aaac 214  
 272 <210> SEQ ID NO: 22  
 273 <211> LENGTH: 184  
 274 <212> TYPE: DNA  
 275 <213> ORGANISM: Homo sapiens  
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 278 gacagcgacg ccgcgagcca gaggatggag ccgcgggcrc cgtggatagr gcaggagrrk 60  
 280 cckgagttt gggacdgra sacasggmaw rtgaaggccc astcastcac agactgaccg 120  
 282 agagaacctg cggatcgcbc tccgctacta caaccagagc gaggccggtg agtgaccccg 180  
 284 gcc  
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296 ccaatactcc ggccctcct gctctatcca cggcgccgc gg	42
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305 <223> OTHER INFORMATION: synthetic.	
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312 <211> LENGTH: 31	
313 <212> TYPE: DNA	
314 <213> ORGANISM: Artificial Sequence	
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323 <210> SEQ ID NO: 26	
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325 <212> TYPE: DNA	
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331 gcgacgtggg ggagtaccgg gcgggtacagg agctggggcg gcctagcgcc gagtaactgga	120
333 acagccagaa ggachtcctg garsrsrgc gsgscsvggt ggacacctac tgcagacaca	180
335 actacggggt tg	192
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339 <211> LENGTH: 23	
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343 <220> FEATURE:	
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**VERIFICATION SUMMARY**

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